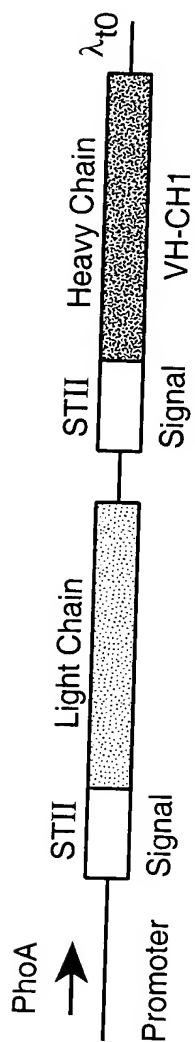


Fab Expression Vector pAK19



Full Length Antibody Expression Vector Derived from pAK19

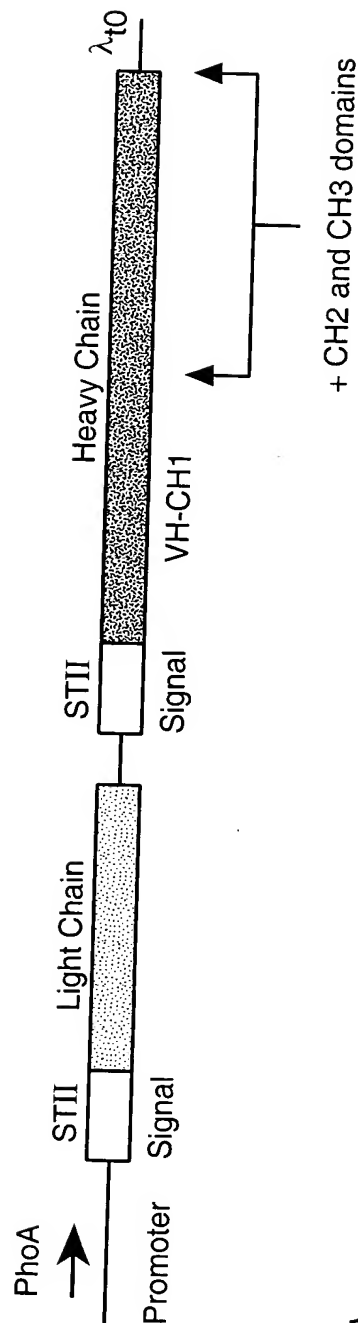
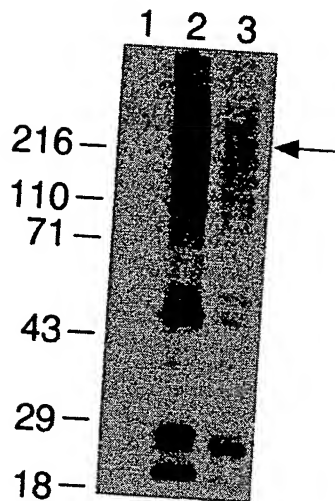
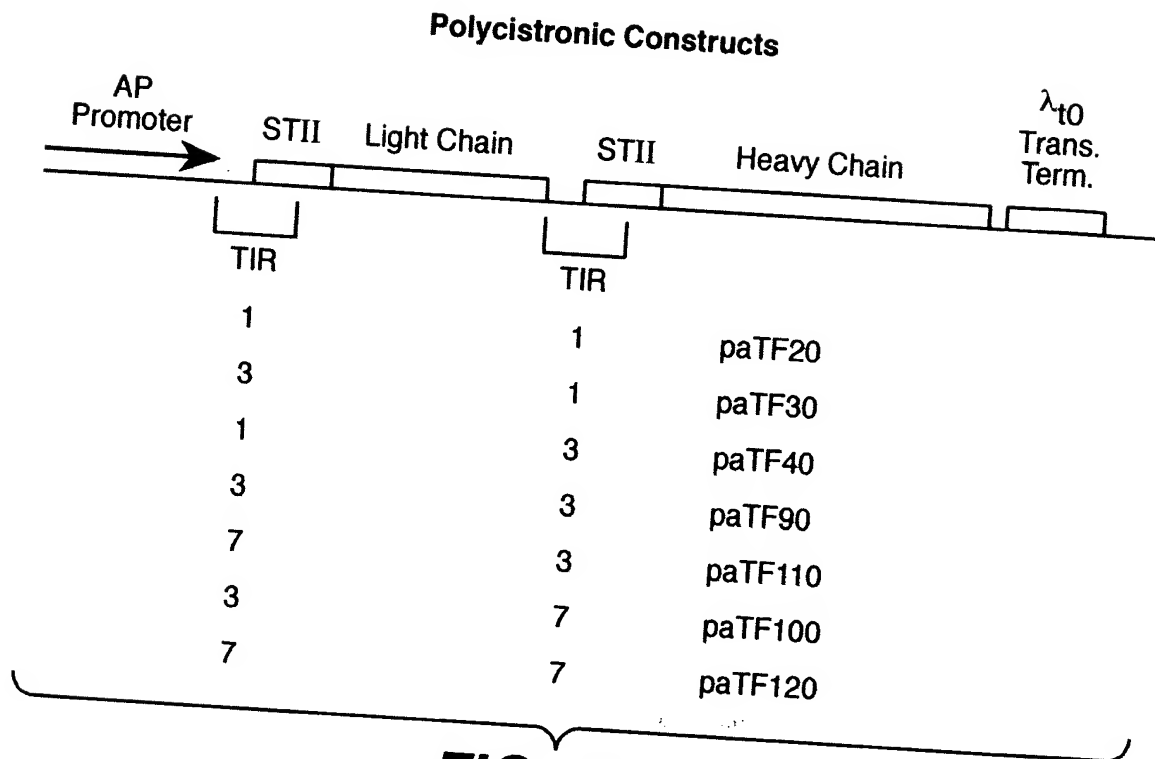


FIG. 1

2/21



**FIG. 2**



**FIG. 3**

3 / 21

FIG.\_4A

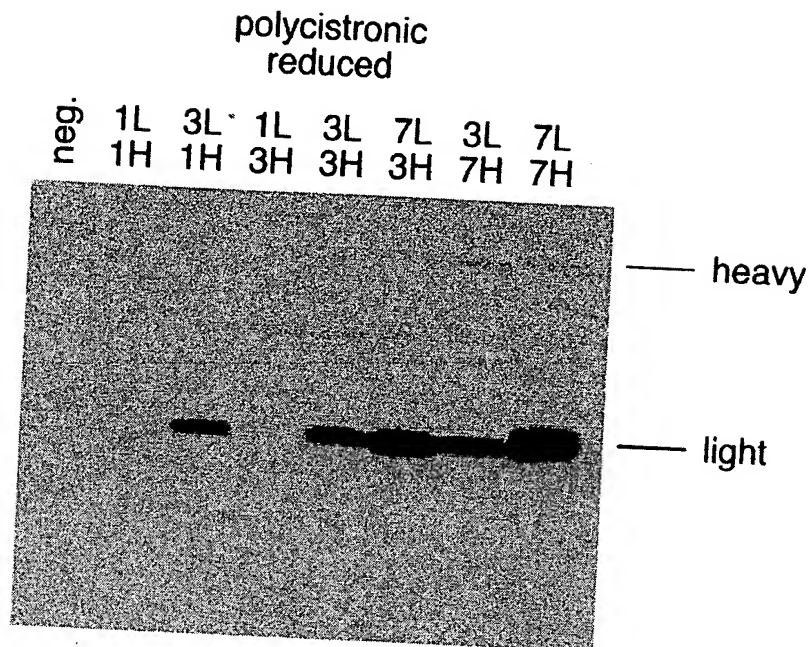
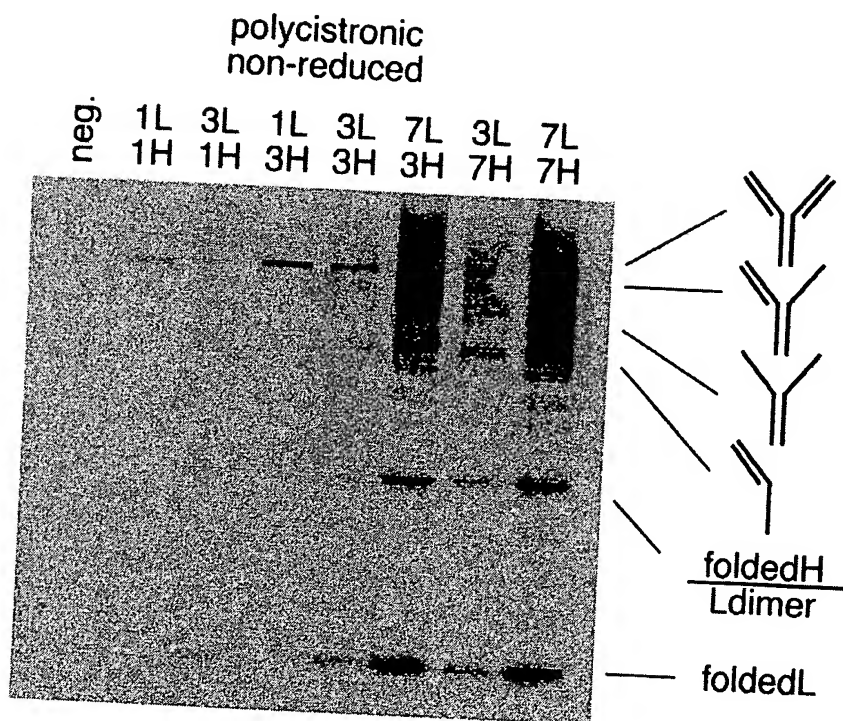
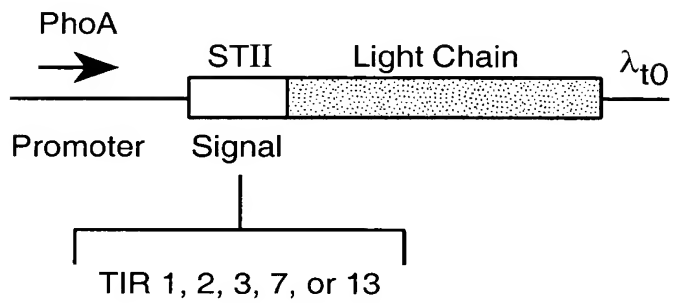
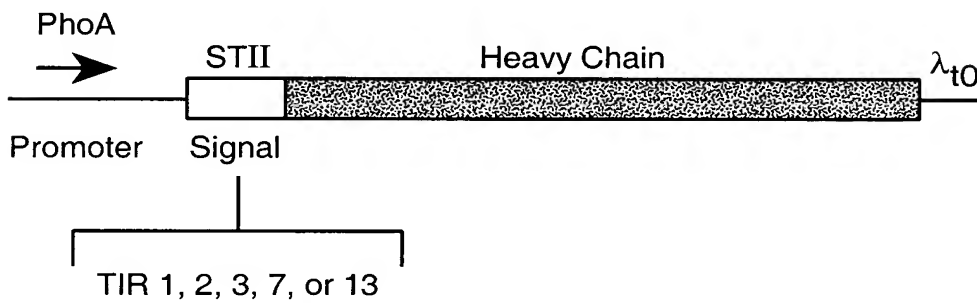
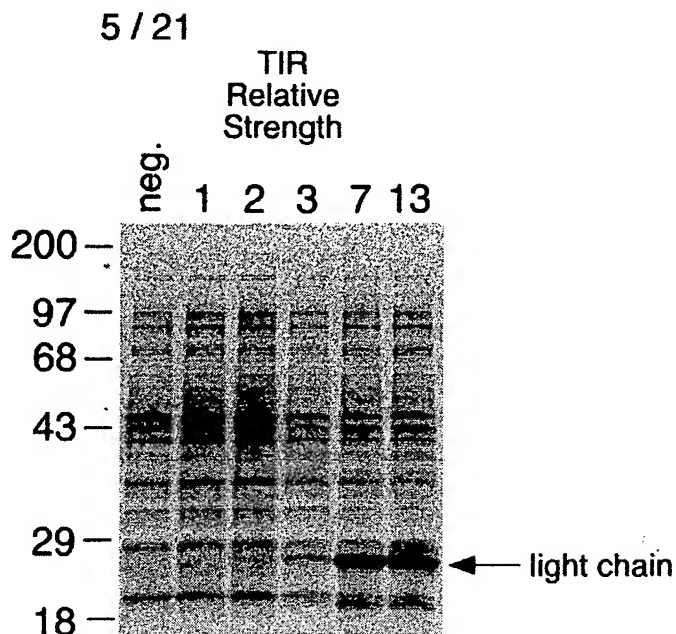


FIG.\_4B

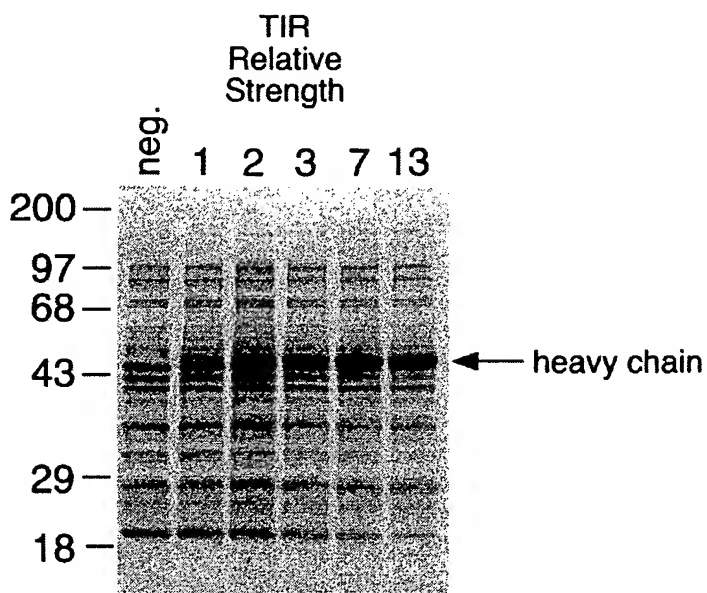


**Light Chain Constructions****Heavy Chain Constructions****FIG. 5**

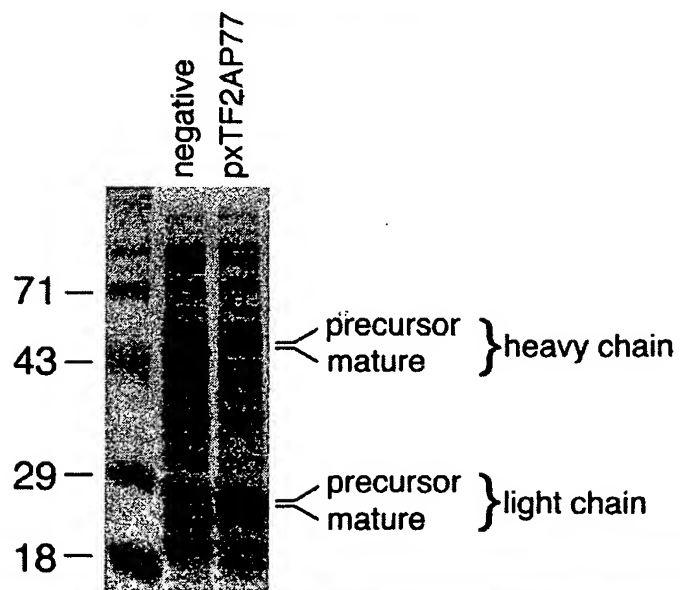
**FIG.\_6A**

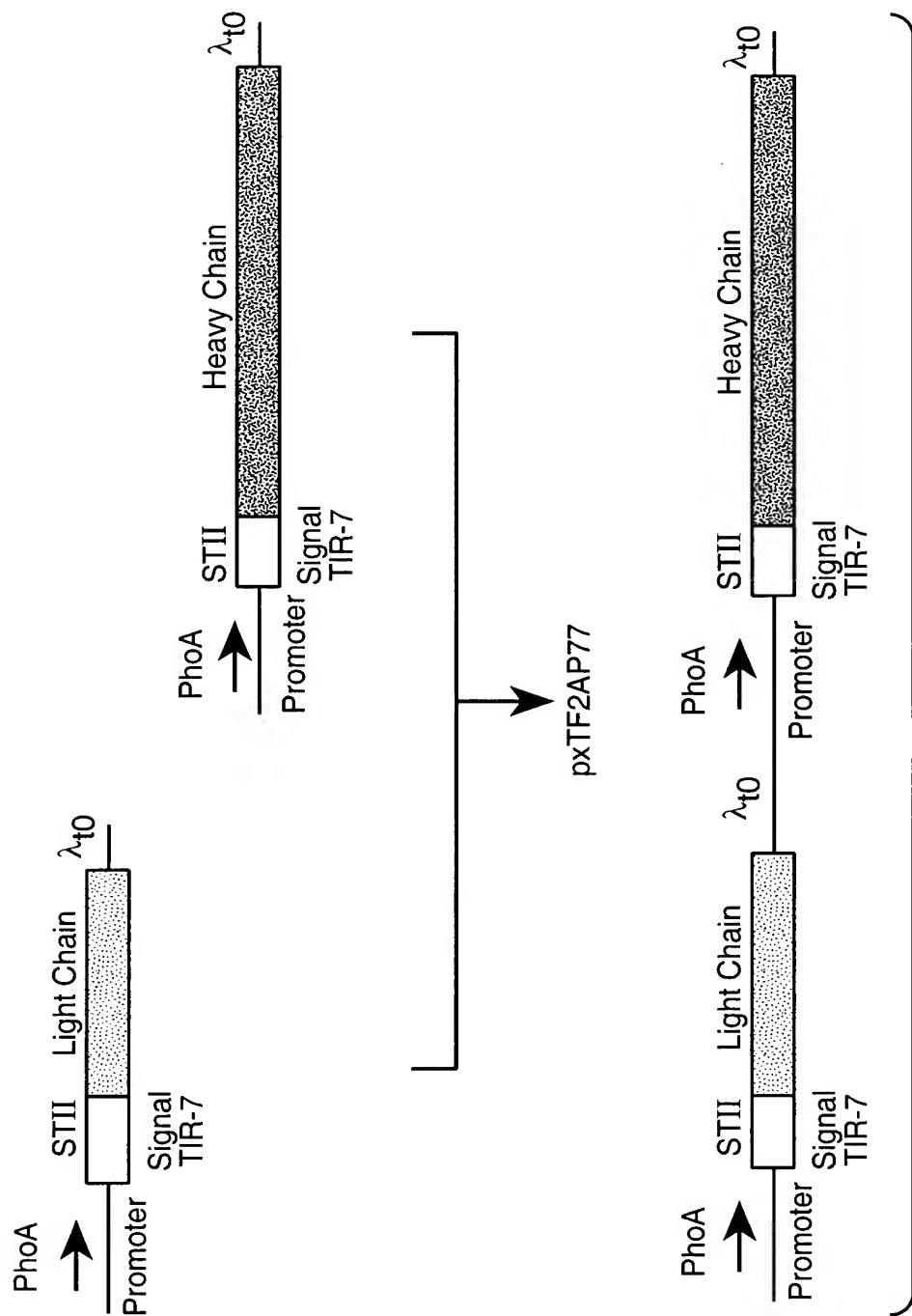


**FIG.\_6B**



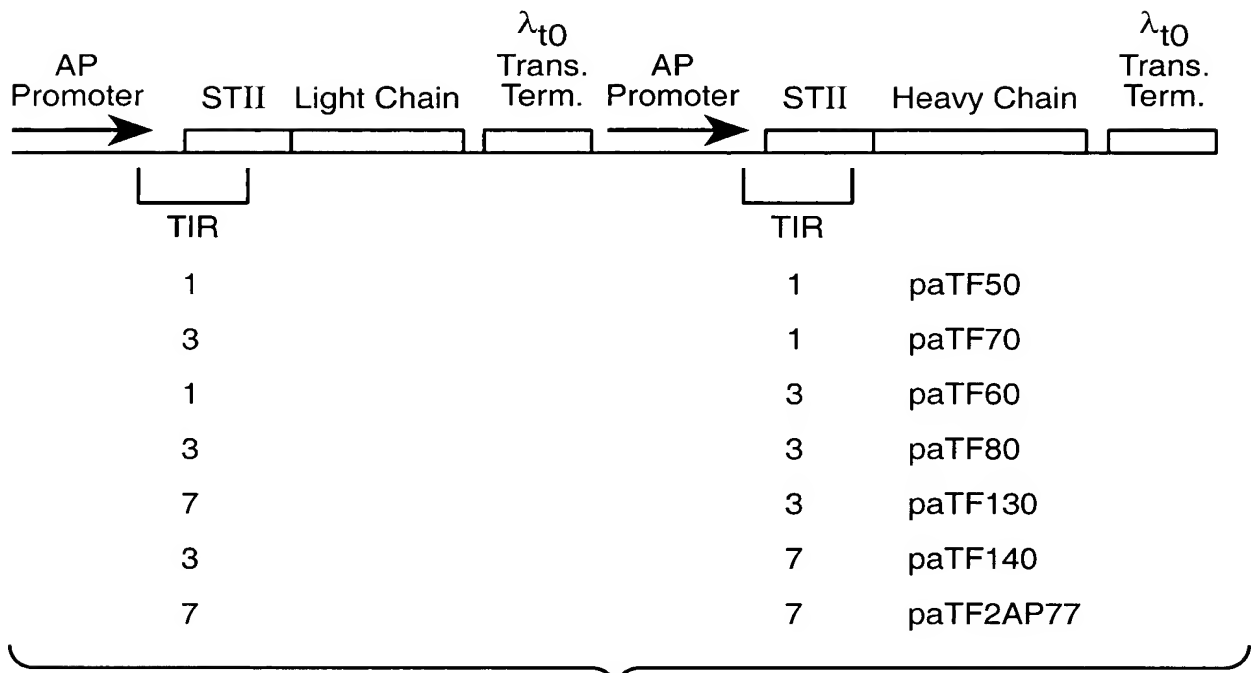
**FIG.\_8**





**FIG.\_7**

### Separate Cistron Constructs

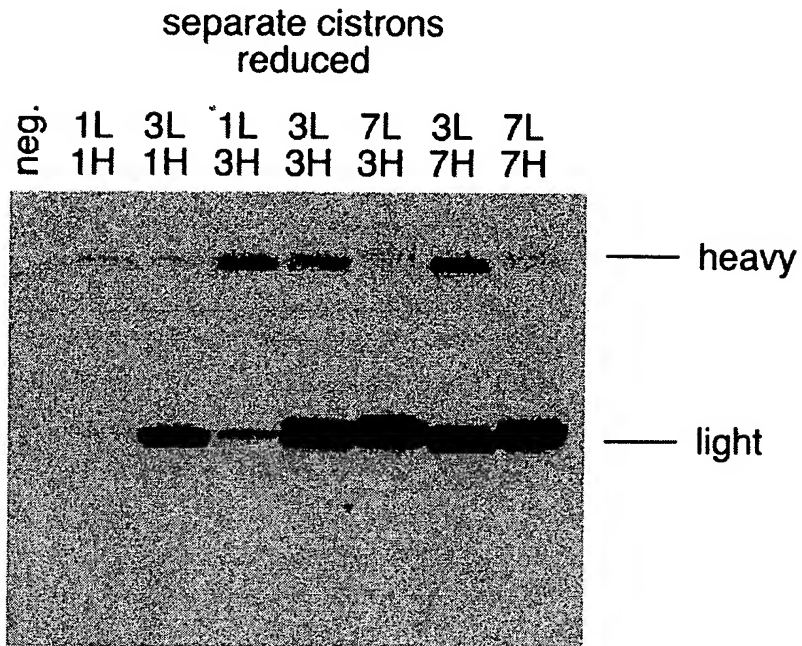


**FIG.\_9**

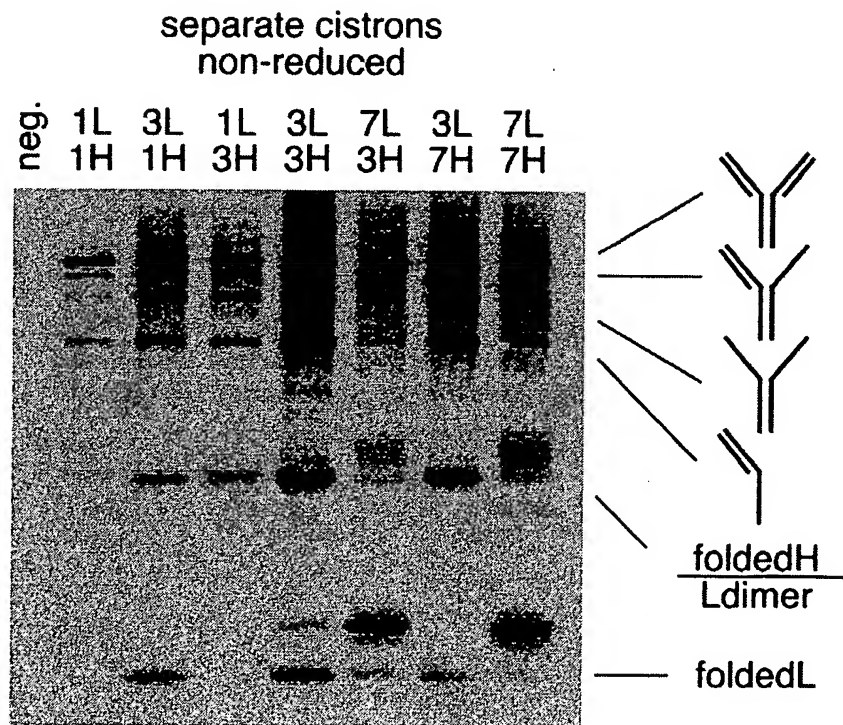
2020-08-20 10:20:00

8/21

**FIG.\_10A**

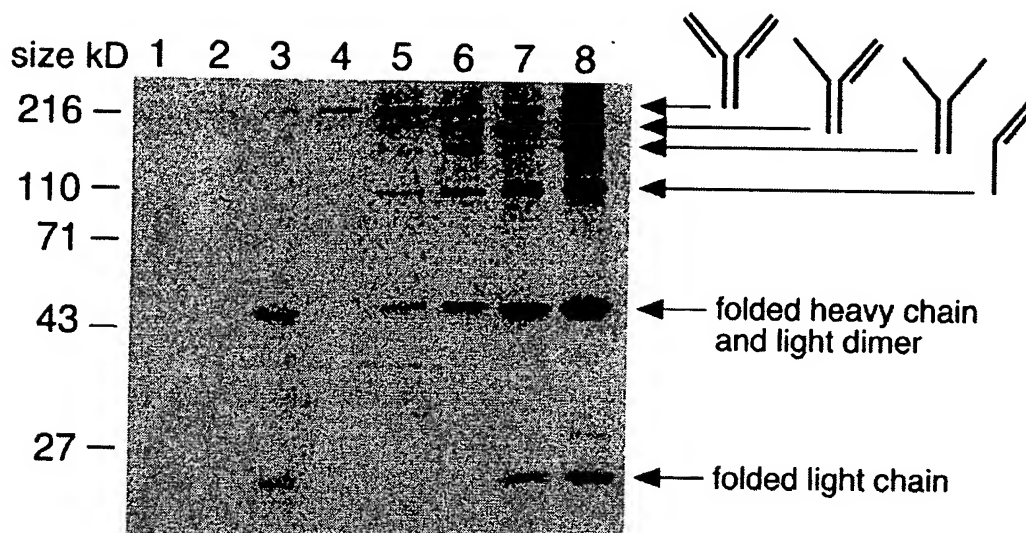


**FIG.\_10B**

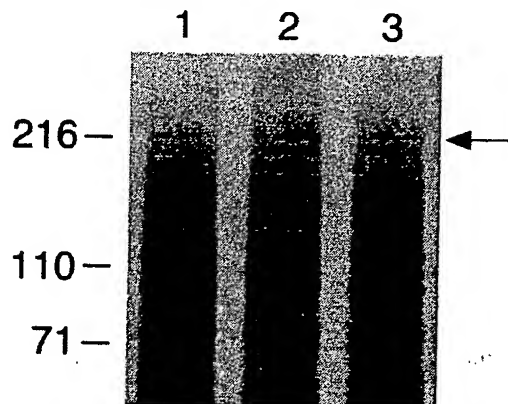




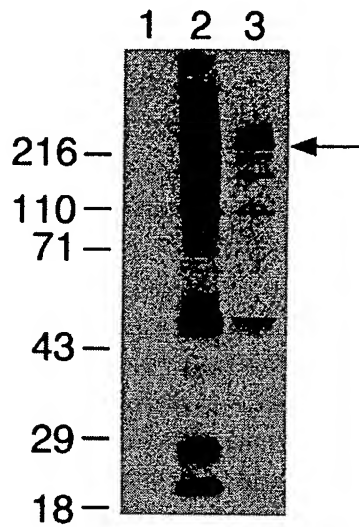
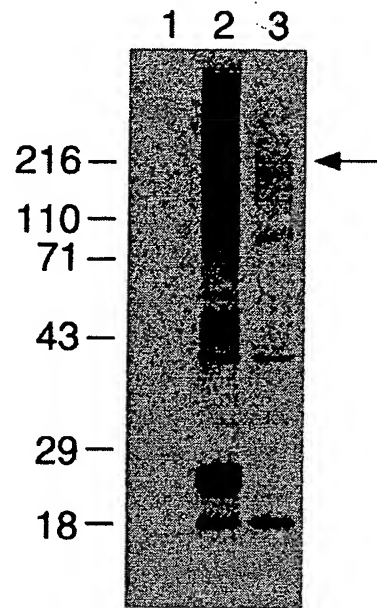
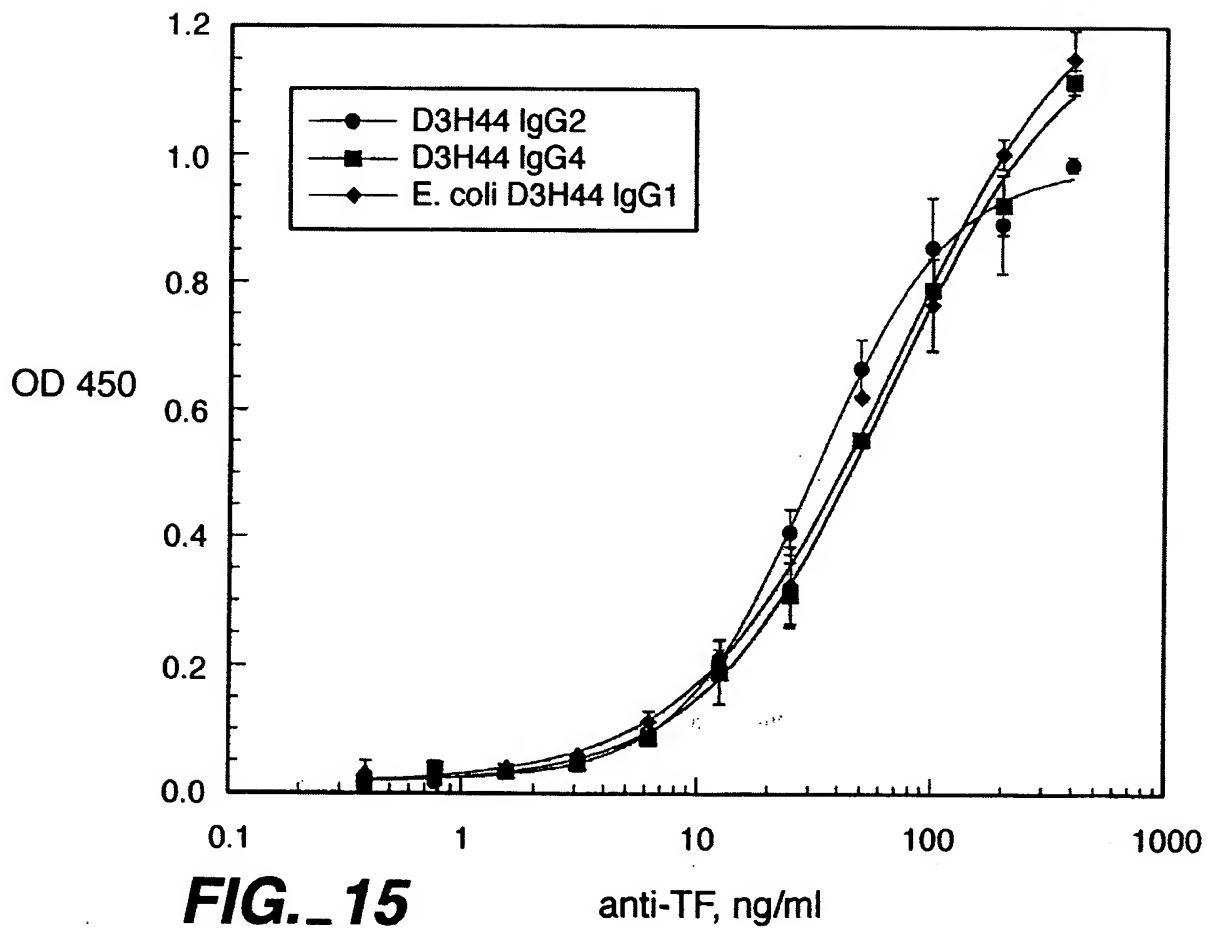
9/21



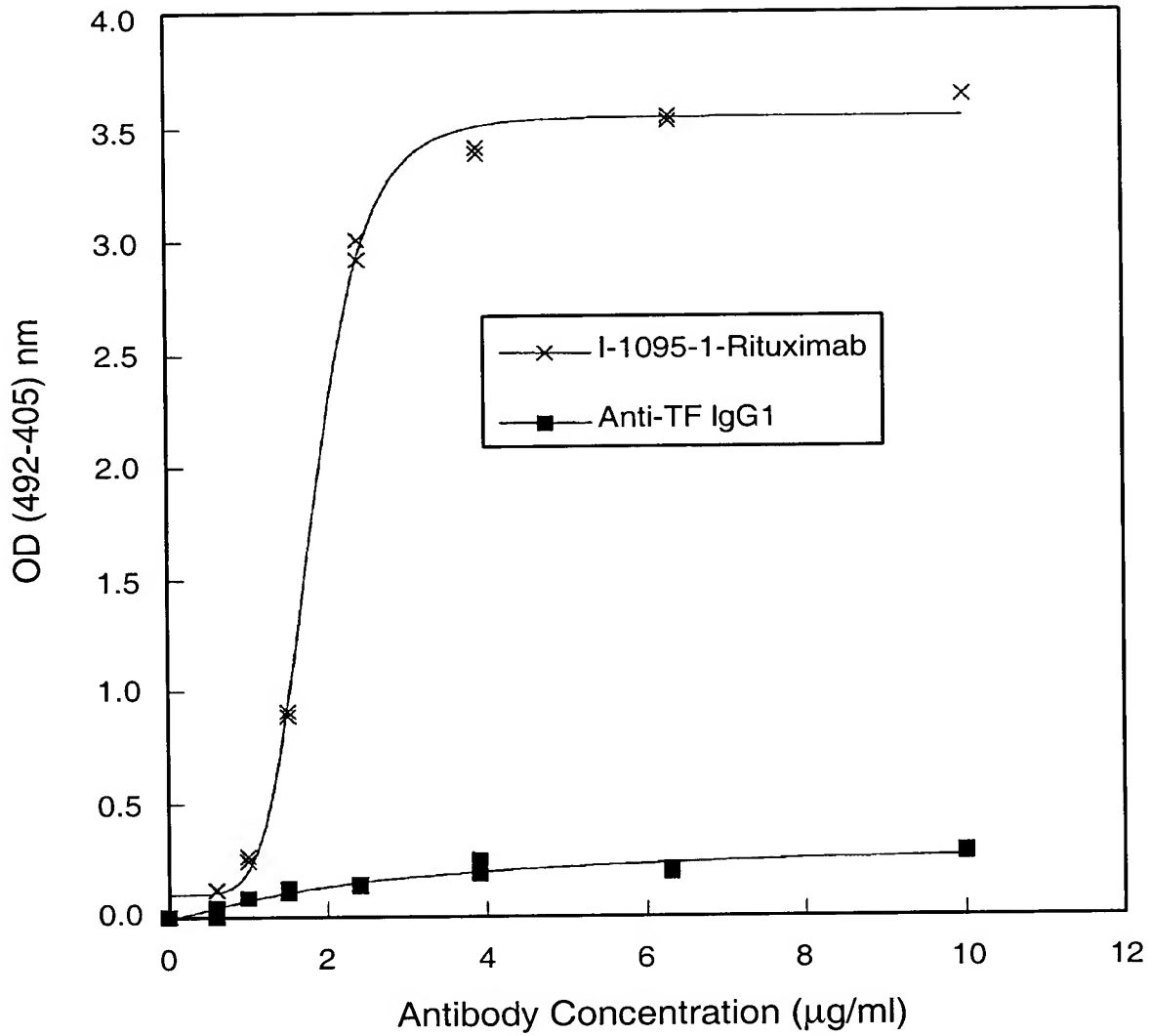
- 1) negative control
- 2) TIR 1-light, TIR 1-heavy, polycistronic
- 3) TIR 3-light, TIR 1-heavy, polycistronic
- 4) TIR 1-light, TIR 3-heavy, polycistronic
- 5) TIR 1-light, TIR 1-heavy, separate cistrons
- 6) TIR 1-light, TIR 3-heavy, separate cistrons
- 7) TIR 3-light, TIR 1-heavy, separate cistrons
- 8) TIR 3-light, TIR 3-heavy, separate cistrons

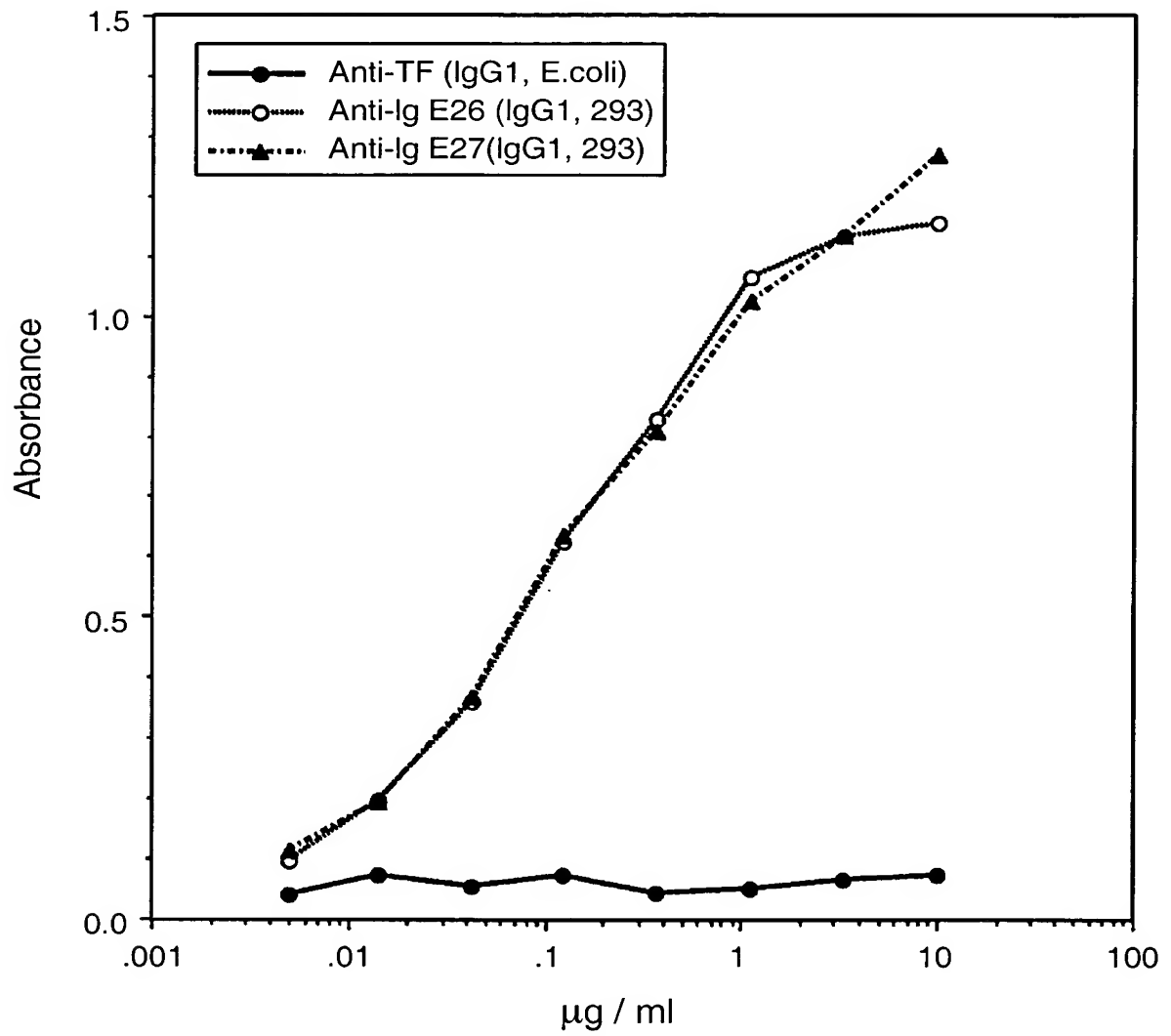
**FIG. 11****FIG. 12**

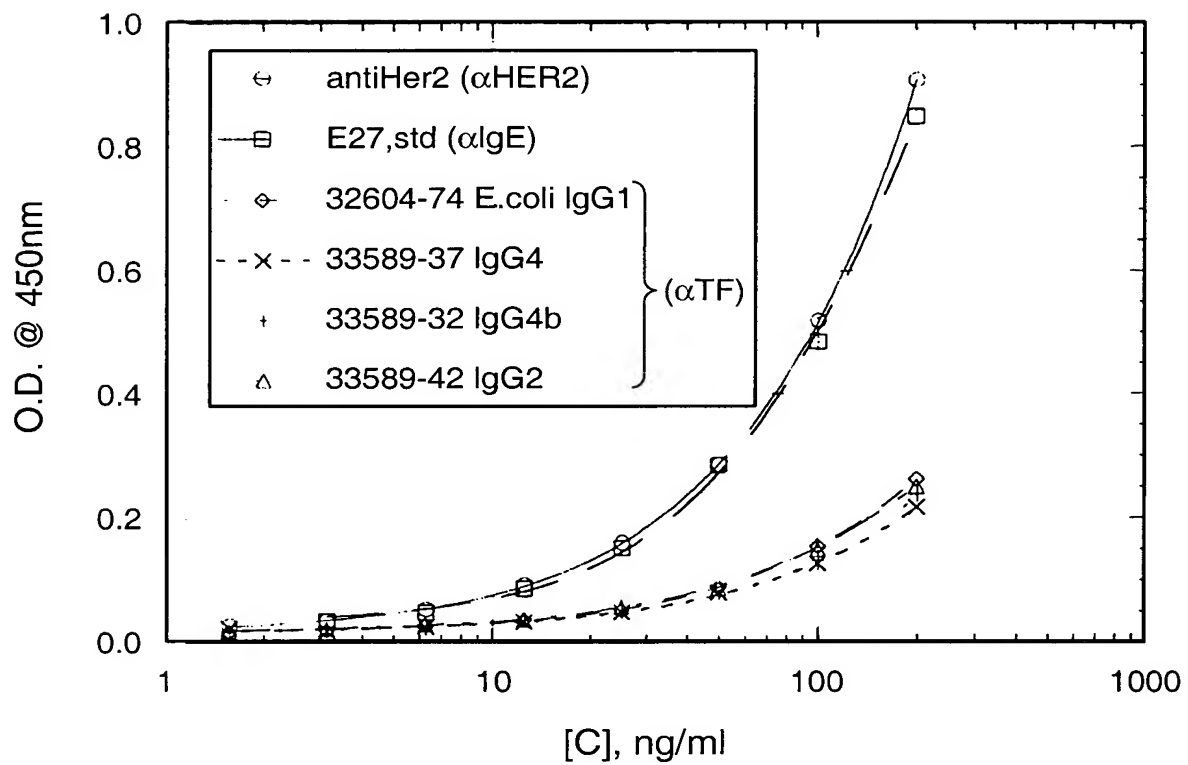
10/21

**FIG. 13****FIG. 14****FIG. 15**

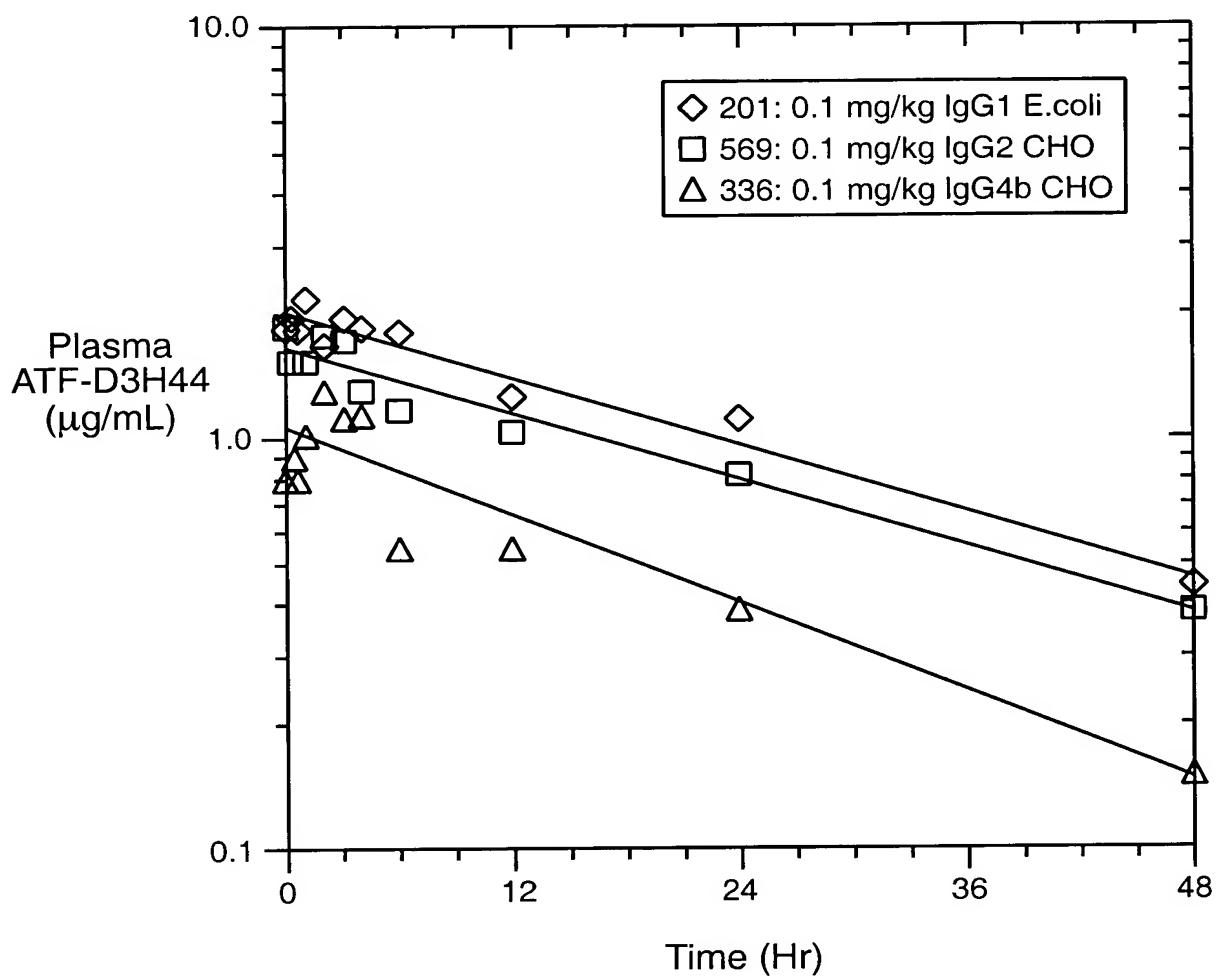
11 / 21

**FIG.\_16**

**FIG.\_17**

**FIG. 18**

14 / 21

**FIG.\_19**

1	GAATTCAACT	TCCTCCATCT	TTGGATAAGG	AAATACAGAC	ATGAAAAATC	TCATGTGCTGA	GTTGTTATTT	AAGCTTGCCC	AAAAAGAAGA	AGAGTCGAAT
	CTTAAGTTGA	AGAGGTATGA	AACTTATTC	TTTATGTCTG	TACTTTTATG	AGTAACGACT	CAACAATAAA	TTGGAACGGG	TTTTTCTTCT	TCTCAGCTTGA
101	GAAGTGTGTG	CGCAGGTAGA	AGCTTTGGAG	ATTATCGTCA	CTGCAATGCT	TCCCAATATG	GCGCAAAATG	ACCAACAGCG	GTTGATTGAT	CAGGTAGAGG
	CTTGACACAC	GCGTCCATCT	TCGAAACCTC	TAATAGCAGT	GACGTTACGA	AGCGTTATAC	CGCGTTTATC	TGGTTGTGCG	CAACTAACTA	GTCCATCTCC
201	GGCGCTGTGA	CGAGGTAAAG	CCCGATGCCA	GCATTCTCTGA	CGACGATPAG	GAGCTGCTGC	GCGATTACGT	AAAGAAGTTA	TTGAAGCATC	CTCGTCAGTA
	CCCGCGACAT	GCTCCATTTC	GGGTACGGT	CGTAAGGACT	GCTGCTATGC	CTCGACGAG	CGCTAATGCA	TTTCTTCAAT	AACTTGTAG	GAGCAGTCAT
301	AAAAGTAAAT	CTTTTCAACA	GCTGTCAATA	AGTTGTCAAG	GCGCAGACTT	ATAGTCGCTT	TGTTTTTATT	TTTTTAATGA	TTTGTAACTA	GTACGCCAAGT
	TTTTCAATTA	GAATAAGTTGT	CGACAGTATT	TCAACAGTGC	CGGCCTCTGAA	TATCAGCGAA	ACAAAAATAA	AAAATTACAT	AAACATTGAT	CATGCGTTCA
401	TCACGTAAAA	AGGGTATCTA	GAATTATGAA	GAAGAATATC	GCATTCTCTC	TTCATCTAT	GTTCTGTTTT	TCATTGTCTA	CAAAACGGTA	CGCTGATATC
	AGTGCATTTT	TCCCATAGAT	CTTAATACTT	CTTCTTATAG	CGTAAAGAAG	AACGTAGATA	CAAGCAAAAA	AGATAACGAT	GTTTGGCAT	GCGACTATAG
1			M K K N I A F L L A S M F V F S I A T N A Y A D I							
			Anti-Tissue Factor Light Chain^							
501	CAGATGACCC	AGTCCCGGAG	CTCCCTGTCC	GCTCTGTGG	GCGATAGGGT	CACCATCAC	TGCAGAGCCA	GTCCGACAT	CAAGAGCTAT	CTGAACCTGT
	GTCTACTGGG	TCAGGGGCTC	GAGGACAGG	CGGAGACACC	CGCTATCCCA	GTGGTAGTGG	ACGTCTCGGT	CAGCGCTGTA	GTCTCTGATA	GACTTGACCA
26	Q M T Q	S P S	S L S	A S V G	D R V	T I T	C R A S	R D I	K S Y	L N W Y
601	ATCAACAGAA	ACCAGGAAA	GCTCCGAAAG	TACTGATTTA	CTATGCTACT	AGTCTCGCTG	AAGGAGTCC	TTCTCGCTTC	TCGTGATCCG	GTCTCTGGAC
	TAGTTGTCTT	TGGTCTCTTT	CGAGGCTTTC	ATGACTTAAT	GATACGATGA	TCAGAGCGAC	TTCTCTCAGG	AAGAGCGAAG	AGACTTAGGC	CAAGACCTGT
60	Q Q K	P G K	A P K V	L I Y Y A T S L A E G V P S R F S G S G S G T						
701	GGATTACACT	CTGACCATCA	GCAGTCTCCA	GCCAGAAGAC	TTCCCAACTT	ATTACTGTCT	TCAGCAOCCA	GAGCTCCAT	GGACATTTGG	ACAGGGTACC
	CTTAATGTGA	GACTGGTAGT	CGTACAGAGT	CGTCTCTG	AAGCGTTGAA	TAATGACAGA	AGTCGTGCT	CTCAGAGGTA	CTGTAAACC	TGTCCCATGG
93	D Y T	L T I S	S L Q	P E D	F A T Y Y C L Q H G E S P W T F G Q G T					
801	AAGTGGAGA	TCAAACGAAC	TGTGGCTGCA	CCATCTGTCT	TCATCTTCCC	GCCATCTGAT	GAGCAGTTGA	AATCTGGAAC	TGCTTCTGTT	GTGTGCTGCT
	TTCCACCTCT	AGTTGTCTG	ACACCGAGT	GGTAGACAGA	AGTAGAAGGG	CGGTAGACTA	CTCGTCAACT	TTAGACCTTG	ACGAAGACAA	CACACGGAGC
126	K V E I	K R T	V A A	P S V F	I F P	P S D	E Q L K S G T A S V V C L L L			
901	TGAATAACTT	CTATCCCAGA	GAGGCCAAAG	TACAGTGGAA	GGTGGATAAC	GCCCTCCAAT	CCGGTAACTC	CCAGGAGAGT	GTACACAGAGC	AGGACAGCAA
	ACTTAATGAA	GATAGGGTCT	CTCCGGTTTC	ATGTCACTT	CCACCTATTG	CCGGAGGTGA	GCCCATTGAG	GGTCCCTCA	CAGTGTCTCG	TGCTGTGCTG
160	N N F	Y P R	E A K V	Q W K	V D N	A L Q S	G N S	Q E S	V T E Q	D S K
1001	GGACAGCAC	TACAGCCTCA	GCAGCACCT	GACGCTGAGC	AAAGCAGACT	ACGAGAAACA	CAAAGTCTAC	GCTGCGAAG	TCACCATCA	GGGCTGAGC
	CTGTGCTGG	ATGTGCGAGT	CGTGGGGA	CTGCGACTCG	TTTGTGCTGA	TGCTCTTTGT	GTTTCAGATG	CGEACGCTTC	AGTGGGTAGT	CCCGGACTCG
193	D S T	Y S L S	S T L	T L S	K A D Y	E K H	K V Y	A C E	V T H Q	G L S
1101	TCGCCCCGCA	CAAAAGACTT	CAACAGGGGA	GAGTGTAAAT	TAAATCTCT	ACGCCGGAAG	CATCGTGGCG	AGCTCGGTAC	CCGGGATCT	AGGCCCTAACG
	AGCGGCGAGT	GTCTTCGAA	GTGTGCCCC	CTCACAATTA	ATTTAGGAGA	TGCGGCCCTGC	GTAGCAOCCG	TOGAGCCATG	GGCCCCTAGA	TCGGGATTCG
226	S P V T	K S F	N R G	E C O						

**FIG. 20a**

1201 CTCGGTTGCC GCGGGGGGTT TTTTATTGTT GCGGACGGC ATCTCGAATG AACCTGCTGC GCAGGTAGAA GCTTTGGAGA TTATCTGCAC TGCATGCTT  
GAGCCAACGG CCGCGCGGCA AAAATAACAA TAGAGCTTAC TTGACACACG CGTCCATCTT CGAAACCTCT AATAGCAGTG ACGTTACGAA

1301 CGCAATATGG CGCAAAATGA CCAACAGGG TTGATTGATC AGGTAGAGGG GCGCTGTGAC GAGGTAAAGC CCGATGCCAG CATTCTCTGAC GACGATAAGG  
GGGTATATAC GCGTTTACT GGTGTGTGOC AACTAACTAG TCCATCTCC CCGGACATG CTCCATTTGG GGCTACGGTC GTAAAGGACTG CTGCTATGOC

1401 AGCTGCTGG CGATTACGTA AAGAAGTTAT TGAAGCATCC TCGTCAGTAA AAAGTTATC TTTCACACAG CTGTATATAA GTTGTACAGG CCGGACTTAA  
TCGACGACGC GCTAATGCAT TTCTTCAATA ACTTCGTAGG AGCAGTCAAT TTTCAAATTAG AAAAGTTGTC GACAGTATTT CAACAGTGGC GGTCTCTGAAT

1501 TAGTGGCTTT GTTTTATTAT TTTAATGTAT TTGTAACTAG TAAGCAAGTT CACGTAAAAA GGGTATCTAG AATTATGAAG AAGAATATCG CATTCTCTCT  
ATCAGCGAAA CAAAAATAAA AAATTACATA AACATTGATC ATGGTTCAAA GTGCAATTTT CCATATAGATC TTAATATCTT TTTCTATAGC GTAAAGAAGA  
M K K N I A F L L  
^STII Signal Sequence TIR-1

1601 TGCAATCTAUG TTGCTTTTCT CTATGTCTAC AAACGGGTAC GCTGAGGTTT AGCTGGTGA GTCTGGCGGT GGCTGGTGC AGCCAGGGGG CTCACTCGGT  
ACGTAGATAC AAGCAAAAAA GATAACGATG TTTCGGCATG CGACTCCAAG TCGACCACTT CAGACCGCCA CCGEACCAAG TCGTCCCTCC GAGTGAGGCA

10 A S M F V F S I A T N A Y A E V Q L V E S G G L V Q P G G S L R  
^Anti-Tissue Factor Heavy Chain

1701 TTGTCTGTG CAGCTTCTGG CTTCAATATT AAGGATGACT ACATGCACTG GGTCCGTGAC GCGCGGGTA AGGCGCTGGA ATGGGTGGA TTGATTGATC  
AACAGGACAC GTGCAAGACC GAAGTTATAA TTCTCTATGA TTGATGTGAC CCAGGCAGTC CCGCGCCCAT TCCCGGAOCT TACCCAAOCT AACTACTAG

43 L S C A A S G F N I K E Y Y M H W V R Q A P G K G L E W V G L I D P

1801 CAGAGCAAG CAACACGATC TATGACCGA AGTTCAGGA CCGTGCCACT ATAAGCGCTG ACAATTCCAA AAACACAGCA TACCTGCAGA TGAACAGCCT  
GTCTGTCTCC GTTGTGTCTAG ATACTGGCT TCAAGGTCTT GGCACGGTGA TATTCGGAC TGTAAAGTT TTGTGTGCT ATGGAAGCTT ACTTGTGGA

77 E Q G N T I Y D P K F Q D R A T I S A D N S K N T A Y L Q M N S L

1901 GGTGTCTGAG GACTGTGCG TCTATTATTG TGCTGAGAC ACGCGCGCTT ACTTCGACTA CTGGGGTCAA GGAACCTGG TCAACGCTC CTCGGCTCC  
CGCAGGACTC CTGTGACGGC AGATAATAAC ACGAGCTCTG TGCGGGGAA TGAAGCTGAT GACCCAGTT CCGTGGGACC AGTGGCAGAG GAGCGGAGG

110 R A E D T A V Y Y C A R D T A A Y F D Y W G Q G T L V T V S S A S

2001 ACCAAGGGC CATCGGTCTT CCGCTGGA CCGCTCTCCA AGACCACTC TGGGGGCACA GCGGCGCTGG GCTGCTGCT CAAGGACTAC TTCCCGGAAC  
TGGTCCCGG GTAGCCAGAA GGGGACCGT GGGAGGAGT TCTGTGGAG ACCCGCGTGT CCGCGGACCC CGACGGAOCC GTTCTCTGATG AAGGGCTTG

143 T K G P S V F P L A P S S K S T S G G T A A L G C L V K D Y F P E P

2101 CCGTGAACGT GTCTGTGAAC TCAGGGGCC TGACCAAGG GTGACCAOCC TTCCCGGCTG TCCPACAGTC CTCAGGACTC TACTCCCTCA GCAGGCTGGT  
GCCACTGCCA CAGCACTTG AGTCCGGGG ACTGTGCGC GCACGTGTGG AAGGGCGGAC AGGATGTGAG GAGTCTGAG ATGAGGGAGT CCGTGCACCA

177 V T V S W N S G A L T S G V H T F P A V L Q S S G L Y S L S V V

2201 GACTGTGCC TCTAGCAGCT TGGGCACCCA GACCTACATC TGCAACGTGA ATCACAAGCC CAGCAACACC AAGGTGGACA AGAAAGTTGA GCCCAATCT  
CTGACACGGG AGATCGTGA ACCCGTGGT CTGGATGTAG ACGTTGCACT TAGTGTTCGG GTCTGTTCGG TTCCACCTGT TCTTTCACCT CCGGTTTGA

210 T V P S S S L G T Q T Y I C N V N H K P S N T K V D K K V E P K S

FIG.--20b



17 / 21

^Start Tet Resistance Coding Sequence

**FIG.-20c**

1 GAATTCAACT TCTCCATACT TTGGATAAGG AAATACAGAC ATGAAAAATC TCATTGCTGA GTTGTATTATTT AAGCTTGCCC AAAAAGAAGA AGAGTCGAAT  
CTTAAGTTGA AGAGGTATGA AACCTATTCC TTATATGCTG TACTTTTTTAG AGTACCGACT CAACAATAAA TTTTCTTCTT TCTCAGCTTA  
101 GAACTGTGT CCGAGGTAGA AGCTTTGGAG ATTATCGTCA CTGCAATGCT TCGCAATATG GCGCAAAATG ACCAACAGCG GTTGATTGAT CAGGTAGAG  
CTTGACACAC GCGTCCATCT TCGAAACCTC TAATAGCAGT GAGGTTACGA AGCGTTATAC CCGGTTTTTAC TGGTTGTGCG CAACTAACTA GTCCATCTCC  
201 GGGCGCTGTA CGAGGTAAAG CCCGATGCCA CCAATTCCTGA CGACGATACG GAGCTGCTGC GCGATTACGT AAAGAAAGTTA TTGAAGCATC CTCGTCAAGTA  
CCCGCGACAT GCTCCATTTC GGGCTACGGT CGTAAGGACT GCTGCTATGC CTGACGCGAG CGCTAATGCA TTTTCTTCAAT AACTTCGTAG GAGCAGTCAT  
301 AAAAGTTAAT CTTTTCACAA GCTGTCAATAA AGTTGTCACG GCGGAGACTT ATAGTCGCTT TGTTTTTTATT TTTTAAATGTA TTTGTAACATA GTACGCAAGT  
TTTTCAATTA GAAAAGTTGT CGACAGTATT TCAACAGTGC CCGCTCTGAA TATCAGCGAA ACAAAAATAA AAAATTACAT AAACATTGAT CATGCGTTCA  
401 TCACGTAAA AGGTATCTA GAATTAAGAA GAAGTAATC CTTCTTATAG CGTAAAGAAG AACGTAGATA CAAGCAAAAA AGATAACGAT GTTTGCGCAT GCGACTATAG  
AGTGCATTTT TCCCATAGAT CTTAATACIT M K K N I A F L L A S M F V F S I A T N A Y A D I  
1  
^STII Signal TIR ~1  
Anti-VEGF Light chain^  
501 CAGTTGACCC AGTCCCGGAG CTCCTGTGTC GCCTCTGTGG GCGATAGGGT CACCATCACG TGCAGCGCAA GTCAGGATAT TAGCAACTAT TTAAACTGGT  
GTCAACTGGG TCAGGGGCTC GAGGGACAGG CCGAGACACC CGCTATCCCA GTGCTAGTGG ACCTGCGGTT CAGTCCCTATA ATCGTTGATA AATTGACCA  
26 Q L T Q S P S S L S A S V G D R V T I T C S A S Q D I S N Y L N W Y  
601 ATCAACAGAA ACCAGGAAA GCTCCGAAAG TACTGATTA CTTACCTCC TCTCTCCACT CTGGAGTCCC TTCTCGCTTC TCTGGATCCG GTTCTGGGAC  
TAGTTGCTTT TGGTCTTTT CGAGGCTTTC ATGACTAAAT GAAGTGGAGG AGAGAGGTA GACCTCAGGG AAGAGCGAAG AGACCTAGGC CAAGACCCTG  
60 Q Q K P G K A P K V L I Y F T S S L H S G V P S R F S G S G S G T  
701 GGATTTCACT CTGACCATCA GCAGTCTGCA GCCAGAAGC TTCGCAACTT ATTACTGTCA ACAGTATAGC ACCGTGCCGT GGACGTTTGG ACAGGTATCC  
CCTAAAGTGA GACTGGTAGT CGTCAGACGT CCGTCTTCTG AAGCGTTGAA TAATGACAGT TGTATATCG TGGCAGCGCA CCTGCAAAACC TGTCCCATGG  
93 D F T L T I S S L Q P E D F A T Y Y C Q Q Y S T V P W T F G Q G T  
801 AAGGTGGAGA TCAACCGAAC TGTGGCTGCA CCATCTGTCT TCATCTTCCC GCCATCTGAT GAGCAGTTGA AATCTGGAAC TGTCTCTGTT GTGTGCTGTC  
TTCCACCTCT AGTTTGTGTTG ACACCGACGT GGTAGACAGA AGTAGAAGG I F P P S D E Q L K S G T A S V V C L L L  
126 K V E I K R T V A A P S V F I F P P S D E Q L K S G T A S V V C L L L  
901 TGAATAACTT CTATCCAGA GAGGCCAAAG TACAGTGGAA GGTGGATAAC GCCCTCCAAT CCGGTAACTC CCAGGAGATG GTCACAGAGC AGGACAGCAA  
ACTTATTGAA GATAGGTTCT CTCCGGTTTC ATGTCACCTT CCACCTATTG CCGGAGGTTA GCCCATTTAG GGTCTCTCTCA CAGTGTCTCG TCCTGTCTGTT  
160 N N F Y P R E A K V Q W K V D N A L Q S G N S Q E S V T E Q D S K  
1001 GGACAGCACC TACAGCTCA GCAGCACCTT GACGCTGAGC AAAGCAGACT ACAGAAACA CAAAGTCTAC GCCTGCGAAG TCACCCATCA GGGCTGAGC  
CCTGTCTGTTG ATGTCGAGT CGTCGTGGGA CTGCGACTCG TTTCGTCTGA TGTCTTTTGT GTTTCAGATG CCGACGCTTC AGTGGGTAGT CCCGGACTCG  
193 D S T Y S L S S T L T L S K A D Y E K H K V Y A C E V T H Q G L S  
1101 TCGCCCGTCA CAAAGACTT CAACAGGGGA GAGTGTAAAT TAAATCTCT ACCTCGGAGC CATCTGTGGG AGCTCGGTAC CCGGGGATCT AGGCTAAACG  
AGCGGGCAGT GTTCTCTGAA GTTGTCCCT CTCACAATTA ATTTAGGAGA TCGGGCCTGC GTAGCACCGC TCGAGCCATG GGGCCCTAGA TCCGGATTGC  
226 S P V T K S F N R G E C O

FIG..21a

1201 CTCGGTGTCC GCCGGGCGTT TTTTATTGTT GCCGACGCGC ATCTCGAATG AACTGTGTGC GCAGGTAGAA GCTTTGGAGA TTATCGTCAC TGCAATGCTT  
GAGCCAACGG CGGCCCGCAA AAAATAACAA CGGCTGCGG TAGAGTTTAC TTGACACACG CGTCCATCTT CGAAACCTCT AATAGCAGTG ACGTTACGAA  
1301 CGCAATATGG CGCAAAATGA CCAACAGCGG TTGATTGATC AGGTAGAGGG GCGCTGTAC GAGGTAAAGC CCGATGCCAG CATTCCTGAC GACGATACGG  
GCGTTATACC GCGTTTACT GGTGTCGCC AACTAACTAG TCCATCTCCC CCGGACATG CTCCATTTCG GGTACGGTC GTAAGGACTG CTGCTATGCC  
1401 AGCTGCTGCG CGATTACGTA AAGAAGTTAT TGAAGCATCC TCCTCAGTAA AAGTTAATC TTTTCAACAG CTGTCAATAA GTTGTACACG CCGAGACTTA  
TCGACGACGC GCTAATGCA TTTCTTCAATA ACTTCGTAGG AGCAGTCATT TTTCAATTAG AAAAGTTGTC GACAGTATTT CAACAGTGCC GGCCTCTGAAT  
1501 TAGTCGCTTT GTTTTATTTT TTTAATGTAT TTGTAAGTAT TACGCAAGTT CACGTAAAAA GGGTATCTAG AATTATGAAG AAGAATATCG CATTTCTTCT  
ATCAGCGAAA CAAAAATAAA AAATTACATA AACATTGATC ATGCGTTCAA GTGCATTTT CCCATAGATC TTAATACTTC TTCTTATAGC GTAAAGAAGA  
1 M K K N I A F L L  
^STII Signal TIR-1  
1601 TGCATCTATG TTCGTTTTTT CTATTGCTAC AAACGCGTAC GCTGAGGTTT ACCTGGTGA GTCTGGCGGT GGCCTGGTGC AGCCAGGGG CTCACCTCCGT  
ACGTAGATAC AAGCAAAAAA GATAACGATG TTTGCGCATG CGACTCCAAG TCGACCACTT CAGACCGCCA CCGGACCACG TCGGTCCCCC GAGTGAGGCA  
10 A S M F V F S I A T N A Y A E V Q L V E S G G L V Q P G G S L R  
^Anti-VEGF Heavy Chain  
1701 TTGTCTGTG CAGCTTCTGG CTACGACTTC ACGCACTACG GTATGAAC TGTCGCTCAG GCGCCGGGTA AGGCCTGGA ATGGGTGGA TGGATTAACA  
AACAGGACAC GTCGAAGACC GATGCTGAAG TCGTGATGCG CATACTTGAC CCAGGCACTC CGGGGCCCAT TCCCGGACCT TACCCAACTT ACCTAATGTT  
43 L S C A A S G Y D F T H Y G M N W V R Q A P G K G L E W V G W I N T  
1801 CCTATACCG TGAACCGACC TATGTCGGG ATTTCAAACG TCGTTTCACT TTTTCTTTAG ACACCTCCA AAGCACAGCA TACCTGCAGA TGAACAGCCT  
GGATATGGCC ACTTGGCTGG ATACGACGCC TAAAGTTTGC AGCAAGTGA AAAAGAAATC TGTGGAGTTT TTCGTGTCTGT ATGGACGTCT ACTTGTGCGA  
77 Y T G E P T Y A A D F K R R F T F S L D T S K S T A Y L Q M N S L  
1901 GCGCGTGTG GACACTGCG TCTATTACTG TGCAAAAGTAC CCGTACTATT ACGGCACGAG CCCTGCTTCC AGCACCCTCTT GGGGTCAAGG AACCTTGGTC  
CGCGCGACTC CTGTGACGGC AGATAATGAC ACGTTTCAAT GGCATGATAA TGCCGTGCTC GGTGACCATTA AAGCTGCAGA CCCCAGTTCC TTGGGACCCAG  
110 R A E D T A V Y Y C A K Y P Y Y Y G T S H W Y F D V W G Q G T L V  
2001 ACCGTCTCCT CGGCCTCCAC CAAGGGCCCA TCGGTCTTCC CCGTGGCACC CTCTTCCAAG AGCACCCTCTG GGGGCACAGC GGCCCTGGGC TGCCTGGTCA  
TGGCAGAGGA GCCGGAGGTG GTTCCCGGGT AGCCAGAAGG GGGACCGTGG GAGGAGTTTC TCCTGGAGAC CCCCCTGTCTG CCGGGACCCG ACGGACCCAGT  
143 T V S S A S T K G P S V F P L A P S S K S T S G G T A A L G C L V K  
2101 AGGACTACTT CCCCAGAACG GTGACGGTGT CGTGGAACTC AGGCGCCTG ACCAGCGGG TGCACACCTT CCGGGCTGTC CTACAGTCTT CAGGACTCTA  
TCCTGATGAA GGGGCTTGGC CACTGSCACA GCACCTTGAG TCCGCGGGAC TGGTCCCGC ACGTGTGGA GGGCCGACAG GATGTACAGA GTCTGTGAGAT  
177 D Y F P E P V T V S W N S G A L T S G V H T F P A V L Q S S G L Y  
2201 CTCCCTCAGC AGCGTGGTGA CTGTGCCCTC TAGCAGCTTG GGCACCCAGA CCTACATCTG CAACGTGAT CACAAGCCCA GCAACACCAA GGTGACAAG  
GAGGGAGTCG TCGCACCACT GACACGGGAG ATCGTCGAAC CCGTGGGTCT GGATGTAGAC GTTGCACTTA GTGTTCGGGT CGTTGTGGTT CCACCTGTTT  
210 S L S S V V T V P S S S L G T Q T Y I C N V N H K P S N T K V D K

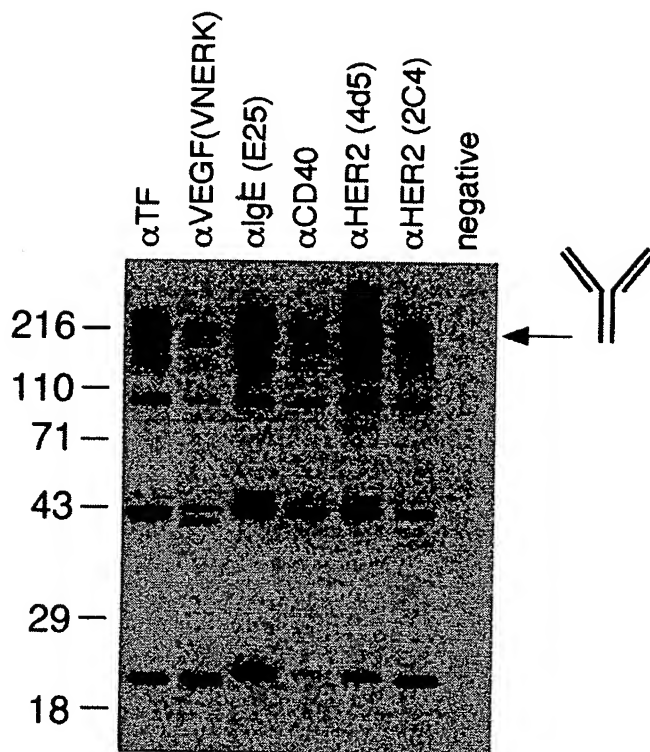
FIG.--21b

2301 AAGTTGAGC CCAAATCTTG TGACAAAACCT CACACATGCC CACCGTGCC AGCACCTGAA CTCCTGGGG GACCGTCAGT CTTCTCTTTC CCCCCAAAC  
 TTTCAACTCG GGTTTAGAAC ACTGTTTGA GTGTGTACGG GTGGCACGG TCGTGGACTT GAGGACCCCC CTGGCAGTCA GAAGGAGAAG GGGGGTTTTG  
 243 K V E P K S C D K T H T C P P C P A P E L L G G P S V F L F P P K P  
 2401 CCAAGGACAC CCTCATGATC TCCCGGACCC CTGAGGTAC ATGGGTGGTG GTGGACGTGA GCCACGAAGA CCTTGAGGTC AAGTTCAACT GGTACGTGGA  
 GGTTCCTGTG GGAGTACTAG AGGGCCTGG GACTCCAGTG TACGCACAC CACCTGCACT CGGTGCTTCT GGGACTCCAG TTCAAGTTGA CCATGCACCT  
 277 K D T L M I S R T P E V T C V V V D V S H E D P E V K F N W Y V D  
 2501 CGGCTGGAG GTGCATAATG CCAAGACAAA GCCGCGGGAG GAGCAGTACA ACAGCACGTA CCGTGTGGTC AGGCTCTCA CCGTCTGCA CCAGGACTGG  
 GCCGACCTC CACGTATTAC GGTTCGTGTT CGGCGCCCTC CTCGTCACTG TGTGCTGCAT GGCACACCAG TCGCAGGAGT GGCAGGACGT GGTCTTGACC  
 310 G V E V H N A K T K P R E E Q Y N S T Y R V V S V L T V L H Q D W  
 2601 CTGAATGGCA AGGAGTACAA GTGCAAGGTC TCCAACAAG CCTCCAGC CCCATCGAG AAAACCATCT CCAAGCCAA AGGCAGCCC CGAGAACCAC  
 GACTTACCGT TCCTCATGTT CACGTTCCAG AGGTGTTTC GGGAGGTCG GGGTAGCTC TTTTGGTAGA GGTTCGGTT TCCCGTCGG GCTCTTGGTG  
 343 L N G K E Y K C K V S N K A L P A P I E K T I S K A K G Q P R E P Q  
 2701 AGGTGTACAC CCTGCCCCA TCCCGGGAAG AGATGACCAA GAACCAAGTC AGCTGACCT GCCTGGTCAA AGGCTTCTAT CCCAGCGACA TCGCCGTGGA  
 TCCACATGTG GGACGGGGT AGGGCCCTTC TCTACTGTTT CTTGTCCAG TCGGACTGGA CGGACCAGTT TCCGAAGATA GGTGCGTGT AGCGGCACCT  
 377 V Y T L P P S R E E M T K N Q V S L T C L V K G F Y P S D I A V E  
 2801 GTGGAGAGC AATGGGCAGC CGGAGAACAA CTACAAGACC AGCCTCCG TGCTGGACTC CGACGGCTCC TTCTTCTCT ACAGCAAGCT CACCGTGGAC  
 CACCTCTCG TTACCCGTCG GCCTCTTGT GATGTTCTGG TGCGGAGGC ACACCTGAG GCTGCCGAGG AAGAAGGAGA TGTCTTCA GTGGCACCTG  
 410 W E S N G Q P E N N Y K T T P P V L D S D G S F F L Y S K L T V D  
 2901 AAGAGCAGGT GGCAGCAGG GAACGCTTTC TCATGCTCCG TGATGCATGA GGCTCTGCAC AACCACTACA CGCAGAAGAG CCTCTCCCTG TCTCCGGGTA  
 TTCTCGTCCA CCGTCGTCCC CTTGCAGAAG AGTACGAGG ACTACGTA CTGAGACGTG TTGGTGATGT GGTCTTCTC GGAGAGGAC AGAGGCCCAT  
 443 K S R W Q Q G N V F S C S V M H E A L H N H Y T Q K S L S L S P G K  
 3001 AATAAGCATG CGACGGCCCT AGAGTCCCTA AGCTCCGTT ACCTCCGTT GTTAACTCAT GTTTGACAGC TTATCATCGA TAAGCTTTAA  
 TTATTCTGTAC GCTGCCGGA TCTCAGGGAT TCGGAGCCAA CGGCGGCCCG CAAAAATAA CAATTGAGTA CAAACTGTCT AATAGTAGCT ATTCGAAATT  
 477 O  
 3101 TCGCGTAGTT TATCACAGTT AATTTGCTAA CGCAGTCAGG CACCGTGTAT GAAATCTAAC AATGCGCTCA TCGTCATCCT CGGCACCGTC ACCCTGGATG  
 AGCCATCAA ATAGTGTCAA TTAAAGATT GCGTCAGTCC GTGGCACATA CTTAGATTG TTACGCGAGT AGCAGTAGGA GCCGTGGCAG TGGGACCTAC  
 3201 CTGTAGGCAT AGGCTTGGTT ATGCCGGTAC TGCCGGGCTT CTTGCGGAT ATCGTCCATT CCGACAGCAT CGCCAGTCAC TATGGCGTGC TGCTAGCGCT  
 GACATCCGTA TCCGAACCAA TACGGCCATG ACGGCCGGA GAACGCCCTA TAGCAGGTAA GGCTGTCTGA GCGTCTAGT ATACCGCAGC ACGATCCGGA  
 3301

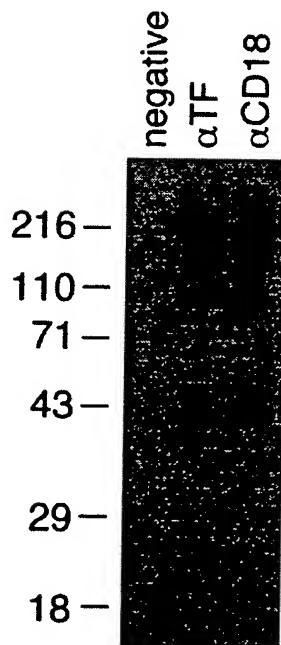
<sup>a</sup>Start Tet Resistance Coding Sequence

FIG.--21c

21 / 21



**FIG.\_22A**



**FIG.\_22B**